40 KNIN K 889 ID NO: 4

RAW SEQUENCE LISTING

DATE: (1/13/20) FATENT APPLICATION: US/09/707,468B TIME: 11:01:10

Input Set : N:\Crf4\12112002\I707468A.raw Output Set: N:\CRF4\01032003\I707468B.raw

```
48 <2115 LENGTH: 19
  49 <212> TYPE: DNA
  50 k213> ORGANISM: Artificial Sequence
  81 k2200 FEATURE:
   52 <223> OTHER INFORMATION: Oligonuplectide primer
   53 <2200 FEATURE:
   84 <2210 NAME/HEY: misc feature
   55 <222> LOCATION: (2)..(2)
   56 <223> OTHER INFORMATION: A or G or C or T/U
   57 (400) SEQUENCE: 4
                                                                                    19
          tnecttgree ecagtarwe
-> 58
   60 <210> SEQ II: NO: 5
   61 <211: LENGTH: 859
   61 <212> TYPE: PRT
  63 (213) ORGANIEM: Mus musculus
   64 <400> SEQUENCE: 5
            Met. Glu Gln Thr Glu Gly Val Ser Thr Glu Cys Ala Lys Ala Ile Lys
                                                 10
   1.1.
            Fro The Asp Gly Lys Ser Val His Gln The Cys Ser Gly Gln Val The
   1.7
                                                                 30
                        20
                                             25
   4-74
            Leu Ser Leu Ser Thr Ala Val Lys Glu Leu Ile Glu Asn Ser Val Asp
   1,4
   110
                                        4()
            Ala Gly Ala Thr Thr Ile Asp Leu Arg Leu Lys Asp Tyr Gly Val Asp
   i
                                    55
                                                         ·50
            Leu Ile Glu Val Ser Asp Asn Gly Cys Gly Val Glu Glu Glu Asn Phe
                                7.0
   4
            Glu Gly Leu Ala Leu Lys His Has Thr Ser Lys Ile Gln Glu Phe Ala
   75
                                                 30
   74,
                            8.5
            Asp Leu Thr Gln Val Glu Thr Phe Gly Phe Arg Gly Glu Ala Leu Ser
   105
   14
                        100
            Ser Leu Cys Ala Leu Ser Asp Val Thr Ile Ser Thr Cys His Gly Ser
   214
                   115
                                        1.20
            Ala Ser Val Gly Thr Arg Leu Val Phe Asp His Asn Gly Lys Ile Thr
   4.
                1.50
                                                         140
            Gir. Lys Thr Pro Tyr Pro Arg Pro Lys Gly Thr Thr Val Ser Val Gin
   83
                                                     155
                                150
   ·- .;
            His Leu Phe Tyr Thr Leu Pro Val Arg Tyr Lys Glu Phe Gln Arg Asn
                                                 110
                            165
   £ 67
            lle Lys Lys Glu Tyr Ser Lys Met Val Glr. Val Leu Gln Ala Tyr Cys
   \dot{\Xi} /
                                             185
                        180
            The The Ser Ala Gly Val Arg Val Ser Dys Thr Ash Gln Leu Gly Gun
                                         200
                                                             205
                    195
   <del>)</del> |
            Gly Lys Arg His Ala Val Val Cys Thr Ser Gly Thr Ser Gly Met Lys
   9]
                                                         220
                                     215
   1)
            Glu Ash Ile Gly Ser Val Phe Gly Gln Lys Gln Leu Gln Ser Leu Ile
   33
                              230
                                                     235
   ٠,٠
            Pro Phe Val Gln Leu Pro Pro Ser Asp Ala Vai Cys Glu Glu Tyr Gly
   35
                                                 250
   زاؤ
            Leu Ser Thr Ser Gly Arg His Lys Thr Fhe Ser Thr Phe Arg Ala Ser
   31
```

RAW SEQUENCE LISTING

HATENT APPLICATI N: US/09/707,468B

DATE: 0.07 -720 03 TIME: 10:00:1

Injut Cet : N:\Crf4\12112002\I707468A.raw
Output Set: N:\CRF4\01032003\I707468B.raw

4 4		2.€€			265			270		
ų į	The His		Arg Thr	Ala Fr	o Gly G	aly Val	Gli. Ali	Thr .	11.5	٠.٠٢
140		275	2		8.Q	•	7 8 5			
181	Ehe Ser	- Ser Ser	Jim Arg	: Gly F	r Wal	Thr III.	Gir Arg	, Gr	1.42	201
162	2.90			291.						
1 1 1	Leu Ser	Met Arg	Fhe Tyt	His M	er Tyr	Ash Arg	His Alm	Tyr	1 1 1	$\mathbb{R}^{k,n}$
104	305		310			315				320
105	Val Val	Leu Asr	. Val Ser	· Val A	sp Ser	Glu Cys	Val Asp	:1e	AST	7.a.l
106			325			330			335	
107 108	The Fro	Asp Lys 340	s Arg Glr. :	. Ile L	eu leu 345	Glm Glu	Olu lys	Leu 350	Leu	Leu
109	Ala Val		Thi Ser	Leu I.	le Glv	Met Fhe	Asp Ser	Āsp	Alā	Asn
110		355			60		365	•		
111	Lvs Let		Asn Gli	. Glm. P.	ro Leu	Leu Asp	Wal Glu	Gly	AST.	Leu
112	370			3115		-	380			
113	Val Lys	. Leu His	Thr Ala	. Gla L	eu Glu	Lys Fro	Val Ero	Gly	Lys	Gli
114	385		340			395				400
115	Asp Asr	. Ser Erc	ser Leo	ı Lys Se	er Thr	Ala Asp	Glu Lys	Arg	ïä.	Ala
110			400			410			415	
117	Ser Ile	e Ser Arç	g Leu Arq	g Glu A		Jer Leu	His Fro		Lys	Glu
118		420			425			430		
119	lie Lys		g Gly Fro			Glu Leu		Ser	i Lite	ero
120		435			40	, T2	445		.	~
1.1			g Gly Val		er ser	Tyr Fro		v či.,	116	ASD
1.2	450		- D Cl-	455	la Aas	Tue Ten	460	11 mas	Thr	Aan
1.3		gry nec	i Arg Gly 470		III Asp	шув њец 475	val per	rio	1111	430
1.4	465 Son Dra	Clu Acr	arc Cys Met		ra Glu		Gla Lvs	Asn	Ser	
1. 6			435			490			495	
1.7 1.8	Leu Ser	Ser Thr 500	: Ser Ala)	a Gly S	er Glu 505	Glu Glu	Fhe Ser	Thr 510	Pro	Glu
1. 9	Val Ala	Ser Ser	Phe Ser	Ser A	sp Tyr	Asn Val	Ser Ser	Leu	Glu	Asp
1:0		515			20		525			
1:1	Arg Fro	Ser Glr	: Glu Thr	r Ile A	sn Cys	Gly Asp	Leu Asp	Суз	Arg	Pro
1 -2	530			535			540			
1:3	Pro Gly	/ Thr Gly	/ Gln Ser		ys Pro		His Gly	Туг	Gln	
1:4	545		550			555				560
1:5 1:6	Lys Ala	i Leu Pro	Leu Ala 565	a Arg L	eu Ser	Pro Thr 570	Asn Ala	ЬУЗ	Arg 575	Fne
137	Lys Thr	Glu Glu	: Arg Fro	Ser A	sn Val	Asn lle	Ser Gln	Arg	Leu	Pro
1.38		580			585			590		
139	Giy Pro	Gln Ser	Thr Ser			Val Asp			Lys	Met
1:0		595			0.0		605			
1;1		-	ı Leu Glr		eu Lys	Ala Gln		Ніз	Glu	Leu
1:2	610			615			620			
1:13		. Lys Glr	: Leu Glr		eu Lys		Asn Lys	HIS	Glu	
1:4	67.5		630		- -	635	and the second	*		640
145	Ser Tyr	Arg Lys	s Fhe Arg	g Ala L	ұз тіе		оту інги	nSH	own one	mada.
146			645			65.0			€55	

RAW SEQUENCE LISTING FATENT APPLICATION: US/09/707,468B DATE: 01/03/2003 TIME: 11:01:10

Input Set : N:\Crf4\12112002\1707468A.raw
Output Set: N:\CRF4\01032003\1707468B.raw

```
Ala Glu Asp Glu Leu Arg Lys Glu Ile Ser Lys Ser Met Fhe Ala Glu
                                            €65
                      660
          Met Glu Ile Leu Gly Gln Fhe Asn Leu Gly Fhe Ile Val Thr Lys Leu
                                                             685
                  675
                                       660
150
          lys Glo Asp Leo Fhe Leo Val Asp Gln His Ala Ala Asp Glo Lys Tyr
                                   695
                                                   Leu Glm Ala Glm Arg Leu
                          leu Gln Gln His Thr Val
          Asn Phe Glu Met
                                                    715
          705
154
          lle Thr Pro Gln Thr Leu Asn leu Thr Ala Val Asn Glu Ala Val leu
155
                                                730
                           725
15.6
          lle Glu Asn Leu Glu Ile Phe Arg Lys Asn Gly Phe Asp Phe Val Ile
15
                                            745
                       740
1: -
          Asp Glu Asp Ala Pro Val Thr Glu Arg Ala Lys Leu Ile Ser Leu Fro
1: +
                                                             765
                  755
                                        760
1( )
          Thr Ser Lys Asn Trp Thr Fhe Gly Pro Gln Asp Ile Asp Glu Leu Ile
10.
                                    775
16...
          Phe Met Leu Ser Asp Ser Pro Gly Val Met Cys Arg Pro Ser Arg Val
1€ ₹
                                                     7.35
                                790
16.
          Arg Gin Met Phe Ala Ser Arg Ala Cys Arg Lys Ser Val Met Ile (ly
1 (
                                                £10
                           805
          Thr Ala leu Asn Ala Ser Glu Met Lys Lys Leu Ile Thr His Met Gly
                                            825
                       820
1 + 3
          Glu Met Asp His Fro Trp Asn Cys Pro His Gly Arg Pro Thr Met Arg
700
                                                             845
                                       840
17
          His Val Ala Ash Leu Asp Val Ile Ser Gln Ash
1:1
                                    855
17.
              850
1/4 <210> SEQ ID NO: €
1. <211> LENGTH: 3056
176 K212> TYPE: DNA
    <213> ORGANISM: Mus musculus
  - <400> SEQUENCE: 6
                                                                                    (S.O.
          gaattooggt qaaqgtootg aagaatttoo agattootga gtatoattgg aggagabaga
1 .
          talcotgtog toaggtaacg atggtgtata tgcaacagaa atgggtgtto ctggagacgc
                                                                                    11.0
1 .
          gtittiticon gagagoggoa obgcaactot obogoggiga oligligacigg aggagioolig
                                                                                    18(1
1 :: 1
          catocatgga goaaacogaa ggogtgagta dagaatgtgo taaggocato aagoctattg
                                                                                    240
          atygyaagts agtopatoaa attigtietg ggeaggigat acteagtita ageaeegetg
                                                                                    200
15 -
          tgaajgagtt gatagaaaat agtgtagatg otggtgotac tactattgat otaaggotta
                                                                                    ( 6 b)
1 - ;
          aaqastatgi igtijiassto ittgaagtti sagasaatig atgtijiggta gaagaigaaa
                                                                                    4.20
1 - 1
          actiting agg totagototing association of obtaining totage to a good action
                                                                                    48.1
1 - 6
          ogsaggttga aactttoggo fittogggggg aagstotgag otstotgtgt goastaagtg
                                                                                    1.....
1: '
                                                                                    (i:)
          atytoactat atotacotgo cacgggtotg caagogttyy gastogacty gtytttgace
1 - :-
          athatgggas satbabbbag saaactebet spoppegabb tasaggased scagtbagtg
                                                                                    651
          tgbagbastt attttatada btaboogtgo gttabaaaga gtttbagagg aabattaaaa
1 : 1
                                                                                    ** 3:1
          aggagtatto baaaatggtg baggtottad aggogtabtg tateatotea geaggogteb
1 1
          gtgtaagetg cactaateag eteggacagg ggaageggea egetgtggtg tgcacaageg
                                                                                    3.1.1
1 ...
                                                                                    3:3:1
          gcacgtotgg catgaaggaa aatatogggt ctgtgtttgg ccagaagcag ttgcaaagcc
7:43
                                                                                    361
           teatteettt igtteagetg ecceetagtg acgetgtgtg tgaagagtae ggestgagsa
1 ...
          cticaggacg scacaaaacs tittctacgi itogggotto atticacagi gcasgcacgg
1 .5
          cgccgqgagg agtgcaacag acaggcagtt tttcttcatc aatcagaggc cctgtgaccc
                                                                                   108)
1 26
```

RAW SEQUENCE LISTING

TATE: 01/03/2013 TIME: 11:01:10 PATENT AFFLICATION: US/09/707,468B

Input Set : N:\Crf4\12112002\I707468A.raw Output Set: N:\CRF4\01032003\I707468B.raw

```
agcasaggto totaagetty tosatgaggt titatesest gistassegg estesgisee
                                                                                1140
19-
          cattiglogt cottaacgti toogtigact cagaatgtgt ggatattaai gtaaciocag
198
          ataaaaggoa aattotasta caagaagaga ngstattgot ggosgiitta aagasotoot
          tgataggaat gittgadagt gatgdaaada (gottaatgt daaddagdag ddaitgdtag
          atgttgaagg taacttagta aagotgcata stgcagaast agaaaagsst gtgccaggaa
          agcaagataa ototoottoa otgaagagoa sagcaqacga gaaaagygta gcatocatot
202
          ccaggotgag agaggoottt totottoato stactaaaga gatcaagtot aggggtocag
                                                                                1500
203
          agactgotga actgacacgg agttttocaa įtgagaaaag gggogtgita tootottato
                                                                                15€
          cttcagacut catctcttac agaggeetes gtggetsgsa ggacaaattg gtgagteeca
                                                                                1623
195
          cggacagoso tggtgactgt atggacagag agaaaataga aaaagactca gggotcagoa
                                                                                1680
          geaceteage tggetetgag gaagagttea geacedeaga agtggeeagt agetttages
                                                                                1740
2.1.7
          gigactatma ogigagotoo otagaagada gadottotoa ggaaaccata aactgiggig
103
          abotogactg cogtostoca ggtacaggac agtocttgaa gccagaagac catggatiato
                                                                                13(1)
1(3
          autgomamy: totacotota getogtetgt pacecacama tgccamagege ttcamagabag
          aggaaaga o otoaaatgto aacatttoto aaagattgoo tggtootoag agcaeotoag
111
          cagotgag: cgatgtagoc ataaaaatga ataagaqaat cgtgctcctc gagttctbtc
                                                                                1040
:12
          tgagttotit agctaagega atgaageagt tacageaect aaaggegeag aacasaestg
113
          aactgagtia cagaaaaitt agggccaaga tttgcchtgg agaaaaccaa gcagbagaag
                                                                                1111
1.14
          atgaacteig aaaagagatt agtaaatega tgtttqmaga gatggagate ttgggtcagt
                                                                                4 6.4
1.15
          ttaacctyg atttatagta accaaactga aagaggacct cttcctggtg gaccagcatg
                                                                               1.28
. 16
                                                                                1340
          ctgcggalja gaagtacaac titgagatgc tgcagcagca cacqgtgctc caggegcaga
. 17
          ggotoatrue accocagact otgaacttaa otgototoaa togaagcigta otgatagaaa
                                                                                4.1
 1.8
                                                                               14+
          atotogasit attoagasag satggottig actitytoat tgatgaggat gotocagtos
. 19
          ctgasagggo tasattgatt toottaccaa ctagtaassa ctggaccttt ggacccaag
...0
          atatagatja actgatotti styttaagtg acagoostgg gytcatgtgc cygosotoac
. 1
          gagtoagada gatgtttjot topagagoot gtoggaagto agtgatgatt ggaabggbgb
                                                                                7:00
          toaatgogag ogagatgaag sagotoatoa opoacatggg tigagatggac cacoobtgga
         actgococca cygoaggoca accatgaggo acgttgocaa totggatgto atototcaga
                                                                                760
. ...:
          actgacadad obcitgtage atagagitta trabagatty troggittge asagagaagg
                                                                                ., €.;;;.
. . 5
         tittaagtaa totgattato gtigtacaaa aattagoaig oigottiaat giaciggalo
. 6
          catttaaasg pagtgttaag goaggoatga tggagtgttb ototagotoa gotaottggg
                                                                                . ) . 1
          tgatocggtg ggageteatg tgageceagg actttgages cacteegage cacatteatg
. .
                                                                                J().):)
          agastcaast saaggacaaa aaaaaaaaga tatttttgaa geettttaaa aaaaaa
                                                                                3056
1-1 <210: SEQ ID NO: 7
1-8 <2110 LENGTH: 862
 -3 <212: TYPE: PFT
1.4 -02130 ORGANISM: Homo sapiens
11-5 <4000 SEQUENCE: 7
          Met Glu Arg Ala Glu Ser Ser Ser Thr Glu Pro Ala Lys Ala Ile Lys
. . . . . 5
                                                                    15
                                                10
           Pro Ile Asp Arg Lys Ser Val His Gln Ile Cys Ser Gly Gln Val Val
.3 53
                                            25
                       20
 3.53
           Leu Ser Leu Ser Thr Ala Val Lys Glu Leu Val Glu Asn Ser Leu Asp
340
                                                            45
                                       40
                   35
 .241
           Ala Gly Ala Thr Asn Ile Asp Leu Lys Leu Lys Asp Tyr Gly Val Asp
 242
                                   55
                                                        60
 243
           Leu lle Glu Val Ser Asp Asn Gly Cys Gly Val Glu Glu Glu Asn Phe
 3 14
                                                                         80
 345
           Glu Gly Leu Thr Leu Lys His His Thr Ser Lys Ile Gln Glu Phe Ala
```

246

RAW SEQUENCE LISTING ERROR SUMMARY FATENT AFFLICATION: US/09/707,468B

TATE: 01/13/2013 TIME: 11:01:11

Input Set : N:\Crf4\12112002\I707468A.raw Output Set: N:\CRF4\01032003\I707468B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seg#:3; N Pos. 5,11 Seq#:4; N Pos. 2

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 4

VERIFICATION SUMMARY

FATENT AFFINCATION: US/09/707,468B

:ATE: 11: -72083 TIME: 11:10:11

Input Set : N:\Crf4\12112002\I707468A.raw
tatput Set: N:\CRF4\01032003\I707468B.raw

D:" M:.VI D: Current Application Number differs, Wrong Format D:48 M:341 W: (46) "n" or "Mad" used, for SEQ ID#:3 after pos.:" D:88 M:341 W: (46) "n" or "Mad" used, for SEQ ID#:4 after pos.:



Creation date: 09-04-2003

Indexing Officer: NNGUYEN7 - NAM NGUYEN

Team: OIPEBackFileIndexing

Dossier: 09707468

Legal Date: 02-24-2003

No.	Doccode	Number of pages
1	SRNT	11

Total number of pages: 1

Remarks:

Order of re-scan issued on